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**ECVPH WORKSHOP, ZURICH 7-9 MAY 2019**

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# ADVANCES IN ADDITIVE BAYESIAN MODEL

## *Outline of the talk*

- ▶ Mixed models – correction for grouped data
- ▶ Heuristic search
- ▶ MCMC over structures
  
- ▶ *Other advanced methods/features:*
  - ▶ *Scoring system*
  - ▶ *Tunable parameter prior*
  - ▶ *Structural prior*
  - ▶ *Data separation*
  - ▶ *Covariate adjustment*
  - ▶ *Likelihood contribution*

## CORRECTION FOR CLUSTERING

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### *Correction for grouped data*

- The way the data were collected has a clear **grouping aspect**
- Then potential for **non-independence** between data points
- Lead to analyses which are **over-optimistic**
- As the **true level of variation** in the data is **under-estimated**
- Could impact study result ... or not!
- **Good practice to check!**

### *In practice:*

- Random effect
- **GLM** -> **GLMM** for each node
- Fit the DAG and check the posterior distribution (**widening**)
- If needed one can **incorporate random effect** in the **scoring** scheme

### *Pitfalls:*

- High computational complexity!

Find maximum a posteriori score

✓ Exact search

▶ Heuristic search

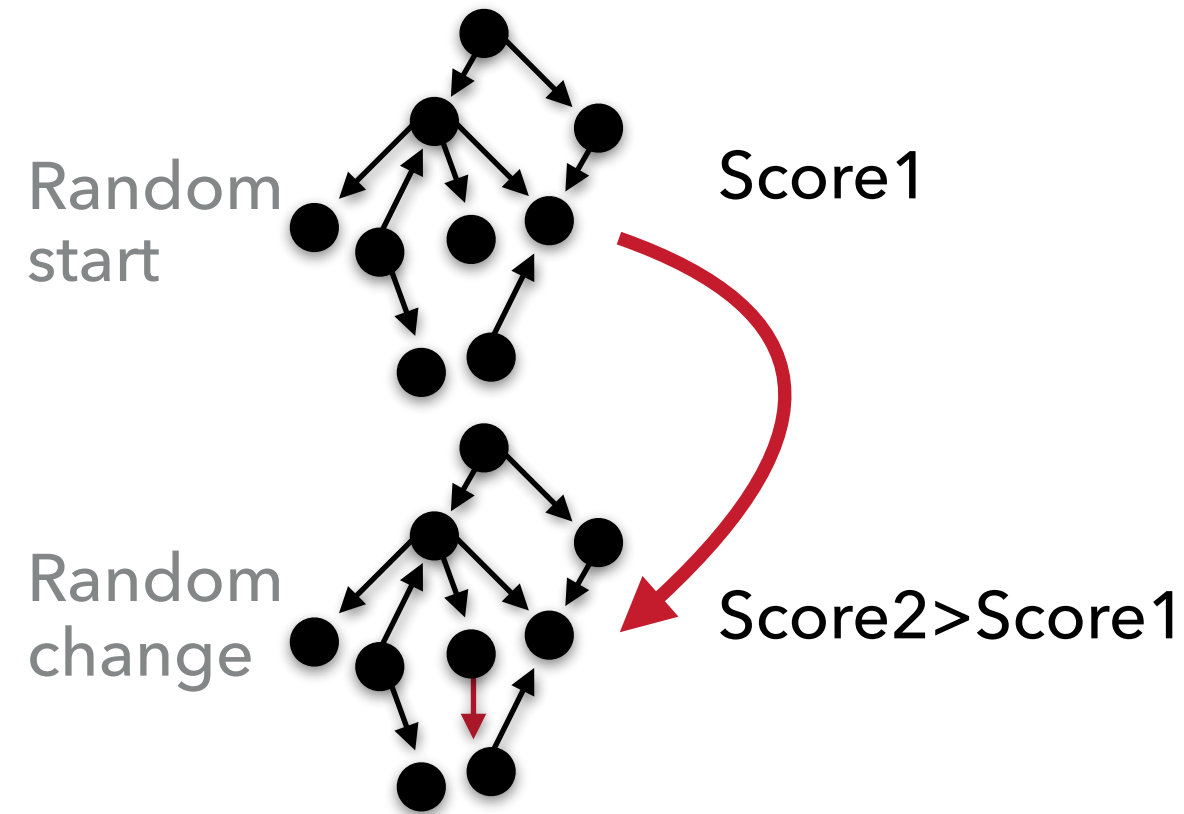
▶ MCMC over structures

# HEURISTIC SEARCH

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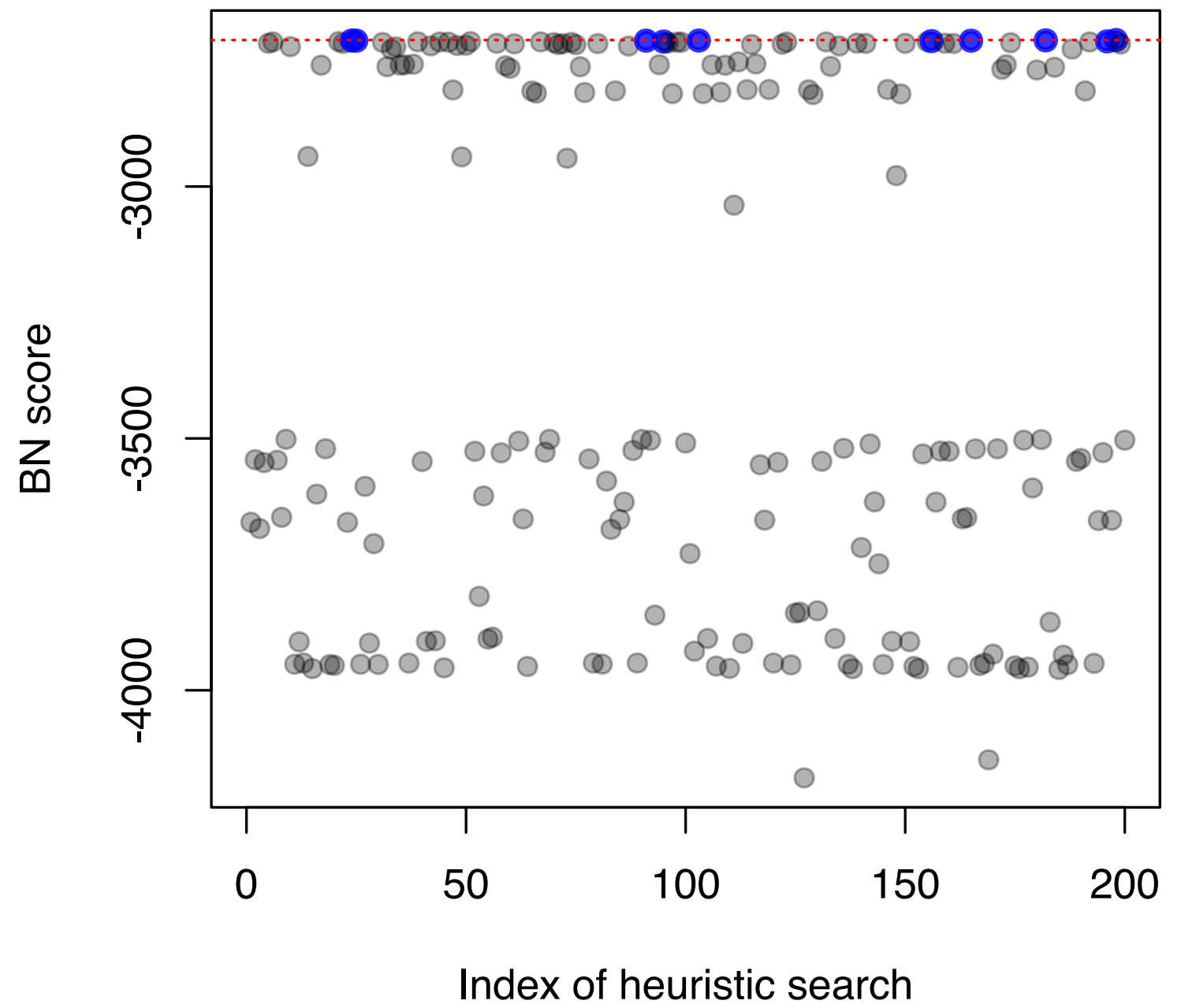
## *Heuristic search: Greedy Hill-Climbing*

- ▶ Simplest heuristic local search
  - ▶ Start with a given network
    - ▶ empty network
    - ▶ best tree
    - ▶ a random network
  - ▶ At each iteration
    - ▶ Evaluate all possible changes
    - ▶ Apply change that leads to best improvement in score
    - ▶ Reiterate
  - ▶ Stop when no modification improves score
- ▶ *Pitfalls:*
  - ▶ Local Maxima
  - ▶ Plateaus
- ▶ *Solution:*
  - ▶ Tabu
  - ▶ Random restart
  - ▶ Simulated annealing

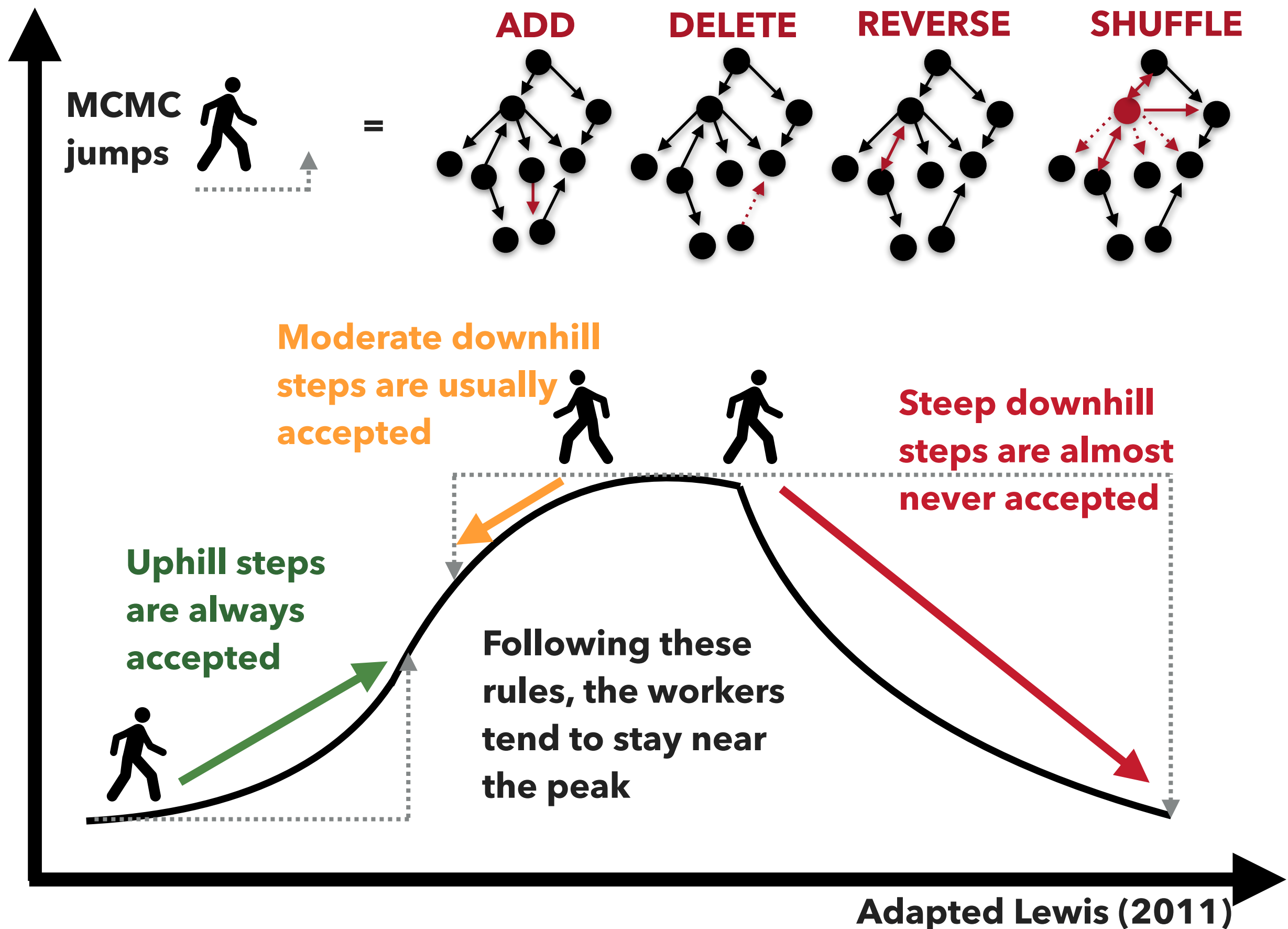


# HEURISTIC SEARCH

```
num.searches <- 200
max.steps <- 150
heur.res <- quiet(search.heuristic(score.cache = mycache,
                                  score = "mlik",
                                  data.dists = dist,
                                  max.parents = 4,
                                  start.dag = "random",
                                  num.searches = num.searches,
                                  max.steps = max.steps,
                                  seed = 3213,
                                  verbose = TRUE,
                                  algo = "hc"))
```

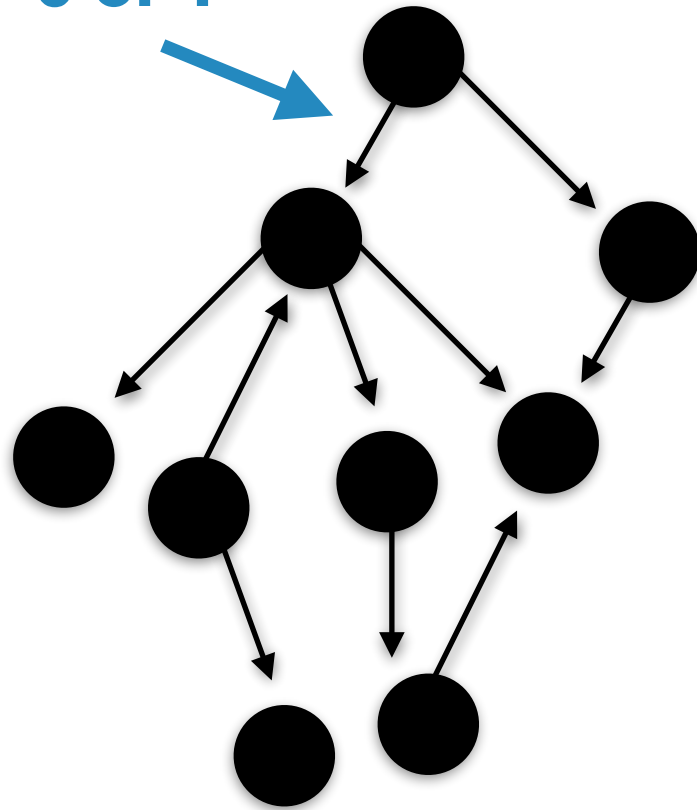


# MCMC OVER STRUCTURES

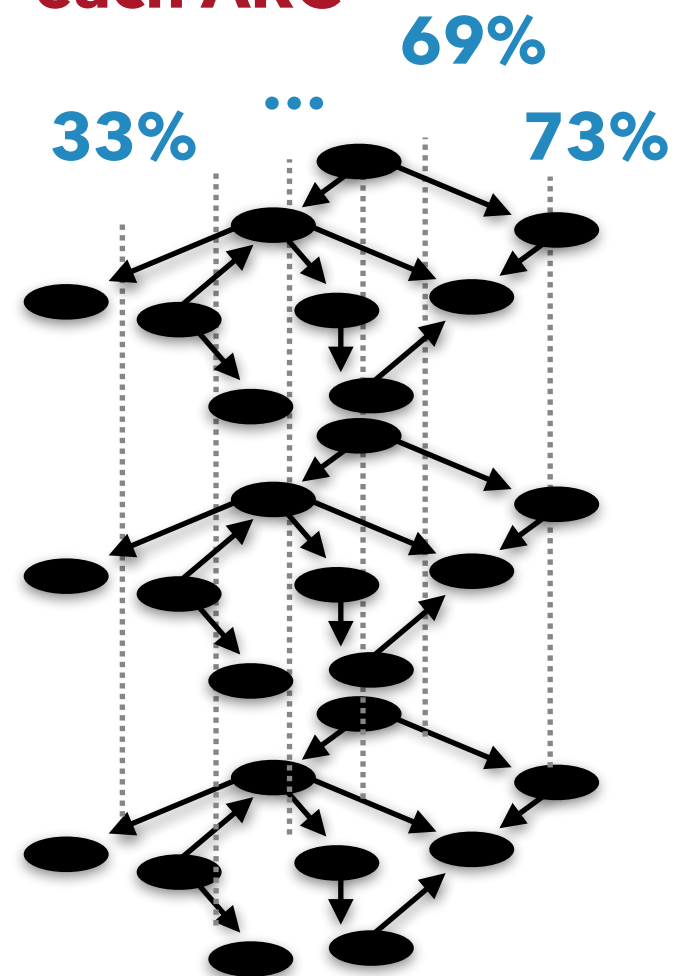


## Best Unique Bayesian Network

0 or 1



## Counting prevalence of each ARC





# MCMC OVER STRUCTURES

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## *MCMC over structures*

- ▶ Selecting the most probable structure
- ▶ Controlling for overfitting
- ▶ Sampling the landscape of high scoring structures
  - ▶ In applied perspective avoid reducing the richness of BN modelling to only **one** structure
  - ▶ Quantify the marginal impact of relationships by marginalising out over structures



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# HANDS-ON EXERCICES

